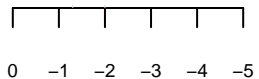
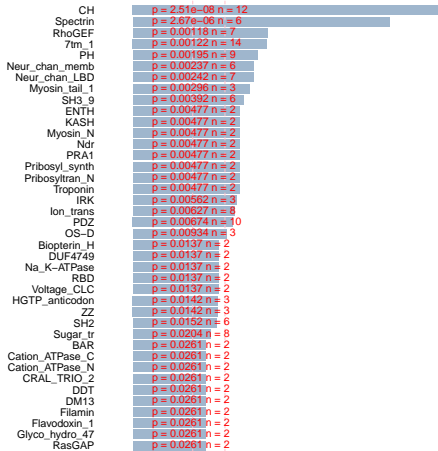


OeCa_M domain



log(p)

n = 752 genes | p adjust BH FALSE